



```

BASE COUNT      303 a      204 c      209 g      232 t      2 others
ORIGIN
9800 Medical Center Drive Rockville, Maryland 20850, USA
Fax : (1) 301 610 8371 Email : fljangel@etech.com URL :
http://nulllength.invitrogen.com

```

Query Match	32.6%;	Score 887.6;	DB 10;	Length 950;
Best Local Similarity	99.98;	Pred. No. 4.5e-134;		
Matches 887; Conservative	1;	Mismatches 0;	Indels 0;	Gaps 0;

Db	311	ctttgsccttcctcctcgcgcctcttgagccggaatccgcggcccgaaaccgcgacttgcga	370
Oy	370	ctttgsccttcctcctcgcgcctcttgagccggaatccgcggcccgaaaccgcgacttgcga	370
Db	1	ctttgcgcgctttccctccgcccgcctttggagccgagatccgcgcccgcgaaacccgcgacttgcga	60
Oy	371	gacggggaactcctactctgctgtagaagccgttagctgaggggaaagagagagggccgtccct	430
Db	61	gacggggaactcctactctgctgtagaagccgttagctgaggggaaagagagagggccgtccct	120
Oy	431	gtcaacagggccggggggaagccgtgtctcttcggagcttcgggtgtgcagacaattcttcggagc	490
Db	121	gtcaacagggccggggggaagccgtgtctcttcggagcttcgggtgtgcagacaattcttcggagc	180
Oy	491	ccagcactgtaagtgccgggagcgtcgcatgtacatcccgagccatgtggaggtccacatgaa	550
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Oy	551	ggacatctccaggttggaatccaagltcaaaaaatgaaaaaaacagaccatctctgaaactc	610
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Db	301	ctgaaacactgtatcaagggccagaaaaatcccaatgtatagccactctggggaaagatattt	360
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Oy	791	gaagctaaatttgcacaaaccttggtgctgaattcttcctgtaccaaagcccgaaatctgcga	850
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Db	601	gacacagctgtctttaaagcagaaggaataattatagttctgtaaaaagctatccaagacaatgat	660
Oy	971	tttaattcccttaaatagtatataataatgtgcttgcattgtaggagtagaaataattccat	1030
Db	661	tttaattcccttaaatagtatataataatgtgcttgcattgtaggagtagaaataattccat	720
Oy	1031	atttgagacatgaatgactcctactcattcttgacaacaaagaaaagaaggtgtatcttaactcaagaa	1090
Db	721	atttgagacatgaatgactcctactcattcttgacaacaaagaaaagaaggtgtatcttaactcaagaa	780
Oy	1091	tcaagtaactctagtaagaatgtggagggcaaaagagttgtatgtgtgtaacaaaanaacaga	1150
Db	781	tcaagtaactctagtaagaatgtggagggcaaaagagttgtatgtgtgtaacaaaanaacaga	840
Oy	1151	acaggaagactcaaaaagcctttgttcaaaagtgagaaagatagaaccga 1198	
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LOCUS	AL581669/c	EST	16-FEB-2001
DEFINITION	AL581669 ltr1_Ft011.Bc1 Homo sapiens cDNA clone CS05d002YE20 3 prime		
ACCESSION	AL581669		
VERSION	AL581669.1		
KEYWORDS	EST.		
SOURCE	human.		
ORGANISM	Homo sapiens		
REFERENCE	Eukaryote; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;		
AUTHORS	Mammalia; Eutheria; Primates; Catarrhini; Homiidae; Homo.		
TTITLE	Li, W.B., Gruber, C., Jesse, J. and Polayes, D.		
JOURNAL	Full-length cDNA libraries and normalization		
COMMENT	Unpublished (2001)		
	Contact: Genoscope		
	Genoscope Centre National de Sequencage		
	Bp 191 91006 Evry cedex - France		
	Email: seqref@genoscope.cns.fr, web : www.genoscope.cns.fr.		
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/sex="male"
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/lab_host="DH10B"
/vector="pCMVSPORT 6; 1st strand cDNA was primed with a NotI-oligo(dT) primer. Five prime end enriched double-stranded cDNA was digested with NotI and cloned into the Not I and Eco RV sites of the pCMVSPORT 6 vector. Library was constructed by Life Technologies. Contact: Feng Liang Life Technologies, a division of Invitrogen, 9800 Medical Center Drive Rockville, Maryland 20850, USA Fax : (1) 301 610 8371 Email : fliang@lifestechn.com URL : http://fulllength.invitrogen.com
```

BASE COUNT	ORIGIN
216 a	146 c 154 g 307 t 22 others

	Query Match	30.0%	Score 817:	DB 10:	length 845:
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DB	845 AMACAACTGMAAGAGACGAGATTTCTGTATAAAGACCCAGGAACTGTAAAAAAGC	786			
QY	1682 tccgtgtatttcacagaccatcccccacccctcaatgtaattgagagggcttaatgga	1741			
DB	785 TCCGKTTTATTTTCAGAGCCATCCCCACCCCTTCMAATGAATWGACAGGCTTAATGGA	726			
QY	1742 aatgagtataataagttccatcgtttaagtaagcgcgaagatagcaataagacgaattta	1801			
DB	725 AAATGACTATTAATTAATCTTCATGTTAAGTACAGCTGAAGATGACATTAAGCAGATTTTA	666			
QY	1802 caacgctacccttacaataaacaacaggaatgcatctctgacatttcggaaccacat	1861			
DB	665 CACAGGTACTCTTACATATAAAACAACAGGAATGATCTTGACATTTCCGAAACACAT	606			
QY	1862 taagtgaaaibgacttaagaagactaaaggtatgactacataaattgaaactaacsagcat	1921			
DB	605 TAAGTGAAATGACTTACAGAACACTAAGGGTGAATCACTAATAATGTAACTACATACAGGAT	546			
QY	1922 ctgtaaatctttctcgt-atttaagtaacagataaagtagatctcaaccaaaaacagagatca	1980			
DB	545 CTGTACATGTTTCTGCACTTTTACGTACAGATTAATAGTGAATGTCACACAAAACGAAAGCA	486			
QY	1981 gatctatgtgctttccacgaacaaagatctcaagaagaaagacctctatctcaatttctt	2040			
DB	485 GATACGTGTGTGTTTTTCCAGCAAAAGATCTTAAGGAAAAGACCTTCATTCATTAATTTTACT	426			

OY 2041 catgattctgctcgtatatacaataaagcttccacaagagccacctactgcttcaaggcaag 2100  
 DB 425 CATGATTCCTGGCTGATATACATTAATAACAGTTTCACAAGACCTACTGTTACAGGCAAG 366  
 OY 2101 gctccattccatactccctccggaaggaacccaatgaaatgtacttcaagaatgtgatagt 2160  
 DB 365 GCCCATTCCTCMTCCTCCCTCCCTSMSCCATGATGTSACTTCAGGAATATGTGATAGT 306  
 OY 2161 ttaaccttcgttaaataatcatatgaataatgaaataatattttagtcgacgaatgaaagaa 2220  
 DB 305 TTMCBCTCTGCTRAATATCATGCAAAAGKRRATTAATTTAGAGAGAAATGAAAAAGAA 246  
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 DB 245 AATCTGGAACCAATGCTGAATTTGATTAAGAACTGAATTTATTACACAAAGAAAAAC 186  
 OY 2281 agaattttagttagtcacgggtacagtccttactagacttgctcagaactagtgaagagaa 2340  
 DB 185 AGAATTTGTGTTACCGGTACAGTCTTTACTAGACTTTGTTCAGACTGTGTAAGAGAAA 126  
 OY 2341 tcaagatttttggtgtttcaagaagctacacagaagaagagtgatattgcaatgttttaagt 2400  
 DB 125 TCAGATTTTGTGGTGTTCACAGCTACACAGAAAGAGTGTATATGCAATGTTTTAGAT 66  
 OY 2401 attcggaagaagaataatcagataatctgttaacagcgcttcttcgtccctccaact 2460  
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 OY 2461 tctac 2465  
 DB 5 TCTAC 1

RESULT 3  
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 LOCUS AL548911 LTI\_NFL006.PL2 Homo sapiens cDNA clone CS0D1043Y010 5  
 DEFINITION prime, mRNA sequence.  
 ACCESSION AL548911  
 VERSION AL548911.1 GI:12884384  
 KEYWORDS EST.  
 SOURCE human.  
 ORGANISM Homo sapiens  
 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 Mammalia; Eutheria; Primates; Catarrhini; Homnidae; Homo.  
 REFERENCE 1 (bases 1 to 769)  
 AUTHORS Li, W.B., Gruber, C., Jessee, J. and Polayes, D.  
 TITLE Full-length cDNA libraries and normalization  
 JOURNAL Unpublished (2001)  
 COMMENT Contact: Genoscope  
 Genoscope - Centre National de Sequencage  
 BP 191 91006 Evry cedex - France  
 Email: seqref@genoscope.cns.fr, Web : www.genoscope.cns.fr.  
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 Location/Qualifiers  
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 /note="Vector: pCMVSPORT 6; Site\_1: NotI; 1st strand cDNA  
 was primed with a NotI-oligo(dT) primer. Five prime end  
 enriched, double-stranded cDNA was digested with Not I and  
 cloned into the Not I and Eco RV sites of the pCMVSPORT 6  
 vector. Library was normalized. Library was constructed by  
 Life Technologies. Contact : Feng Liang Life Technologies,  
 a division of Invitrogen 9800 Medical Center Drive  
 Rockville, Maryland 20850, USA Fax : (1) 301 610 8371  
 Email : fliang@lifestech.com URL :  
 http://fulllength.invitrogen.com"

BASE COUNT 249 a 168 c 170 g 182 t

Query Match 28.3%; Score 769; DB 10; Length 769;  
 Best Local Similarity 100.0%; Pred. No. 6,9e-115;  
 Matches 769; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 339 agccgagatccggcccccgggaaccccgacctgacagcggttacctactcgttagagacc 398  
 DB 1 AGCCGATCCGGCCCGCGGAACCCGACCTGCAGACCGCGGTACTCTACTCGGTAAGAGCC 60  
 OY 399 gtagctggcggaaggaagagcgccgctcgttaacacggccggggggaagccggtctt 458  
 DB 61 GTAGCTGGGGGAAGGAGAGGGGGCGCTCTGTCAACAGCGCGGGGAAGCGGTCTCTTT 120  
 OY 459 ccgctgcccgggtgtagacatttcccgaccacagatgtagtgcggcgagaccga 518  
 DB 121 CCGGCTGCCCGCTGCCGACATTTCTCCGACCCACATGTAGCTCCGGCGACTGCCA 180  
 OY 519 tgaactccggagccatgagagatccacagataagagacattccagggctggatccaagta 578  
 DB 181 TGAACTCCGGAGCCATGAGGATCCAGTAAGGACATTTCCAGGTGGAAATCCAAGTCA 240  
 OY 579 aaaaagaaaaaacagacacatctcgaatctcgaatcgaatcgaacgagccagaanaat 638  
 DB 241 AAAATGAAAAAAACAGACCATCTCTGAAAATCTCTGAAAATGATAACAGCCAGAAAAAT 300  
 OY 639 ccaaatgtaagccactttggggaagaagtatttacctgactactctctgacatat 698  
 DB 301 CCAAATGTAAAGCCACTTGTGGGAAAGTATTTACCTTCACTTACCTCTGTACCATAT 360  
 OY 699 ctgaaaaacttcaaaaggaacatlaagatcgtggagggagctgagaattctcagca 758  
 DB 361 CTGAAAAACTTCAAAAGGACATTAAGATCTGGAGGGGAGTGAAGAAATTTCTAGCA 420  
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 DB 421 AAGATATCAGTTATTTATTATTCAATTAAGAAAGAGACGTAATTTGCAACAACTTGAGTC 480  
 OY 819 gaatttctctgtacccaagtcgaatcgaatcgaatcgaatcgaatcgaatcgaatcga 878  
 DB 481 GAATTTCTCTCTGTACCAAGTCCAGAAATCTGCATATCTGCAGAAACACTCTCATC 540  
 OY 879 ccagccatgatgaagttatatttaagtaacacagacagtggttttaagcagaagaaat 938  
 DB 541 CCAAGCATGATGCAATTTCTTTAATGTCACACACAGTGTGTTTAAGCAGAGGAAAT 600  
 OY 939 tattaagtgaaaaagctatcaagaagacatgatttatttccctcaaatagatatatcaaa 998  
 DB 601 TATTAGTTGAAAAAGCTATCAAGAGACCATGATTTATTCCTCAAAATGATATTTATCA 660  
 OY 999 atgcttgcacagggaggaataaattcttcatatgtagacatgatatcatatgaac 1058  
 DB 661 ATGCCCTTGATCGGGAGGAAAAATTTCTCATATTTATGATGATATCATATCATATTGAAC 720  
 OY 1059 aaaaagaaaaagtgtagtatttactcaagaagaatcaagtgactcagtaag 1107  
 DB 721 AAAAGAAAAAGAGTTGTAATTTACTCAAGAAATCAAGTACTTGATAG 769  
 RESULT 4  
 AL573294 784 bp mRNA EST 16-FEB-2001  
 LOCUS AL573294 LTI\_NFL006.PL2 Homo sapiens cDNA clone CS0D1043Y010 3  
 DEFINITION prime, mRNA sequence.  
 ACCESSION AL573294  
 VERSION AL573294.1 GI:12932397  
 KEYWORDS EST.  
 SOURCE human.  
 ORGANISM Homo sapiens  
 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 Mammalia; Eutheria; Primates; Catarrhini; Homnidae; Homo.  
 REFERENCE 1 (bases 1 to 784)  
 AUTHORS Li, W.B., Gruber, C., Jessee, J. and Polayes, D.  
 TITLE Full-length cDNA libraries and normalization

JOURNAL  
COMMENT unpublished (2001)  
Contact: Genoscope  
Genoscope Centre National de Sequencage  
BP 191 91006 Evry cedex - France  
Email: seqref@genoscope.cns.fr, Web : www.genoscope.cns.fr.

FEATURES  
source Location/Qualifiers

1. 784  
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BASE COUNT 234 a 135 c 126 g 285 t 4 others

ORIGIN

Query Match 28.38; Score 768.2; DB 10; Length 784;  
Best Local Similarity 99.48; Pred. No. 9.2e-115;  
Matches 778; Conservative 3; Mismatches 1; Indels 1; Gaps 1;

1830 ggaatgacatcttcgacatcttcgacacacacatlaagtgaaatgaacttagaagaactaaag 1889  
784 GGAATGACATCTTTGACATCTTCGACACACACATTAAGTAAATGACTTAACAACAACTAAG 725  
1890 ggtatgacacataaataatgatacagagcctgtacatgttttcgatttcgatacaga 1949  
724 GGTATGACACATAAATAATGATAACAGGCACTGTACATGTTCGATTTCGATTCAGTACAGA 665  
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664 TAATAGTGATCTCAACACACACAGAGTGCAGATACGTGCTTTTCCAGCAAGATC 605  
2009 tcaagaaagagacaccttcattcaatatttaccatgatttcggtcgtatatacaataaaca 2068  
604 TCAGAGAAAGAGACCTTCATTCAATATTACTCATGATTCGTGCTGTATACAAATTAACA 545  
2069 gtccacagagacaccttcattcaatatttaccatgatttcggtcgtatatacaataaaca 2128  
544 GTTCACAGAGACACCTTCATTCAATATTACTCATGATTCGTGCTGTATACAAATTAACA 485  
2129 ccaatgaaatggaactcaagaataatgataatgttcccttcgtgtaaaatcatcagaaga 2188  
484 CCNAATGAATGGACTTCAGAAATATGATATGTTTACCTTGTGTAATATCATCGAAGAAG 425  
2189 tgaataataatcttagagacgaataatagaagaagaatctggaacccaatgctgaaatcga 2248  
424 TGAATAATATATATATGAGACGAATATAGAAAGAAATCTGGAACCAAAATCGCAATTTGATA 365  
2249 aaagaacggaattattacacagaagaagaagaatgttgaagtcacacggtacagtcct 2308  
364 AAAGAACGGAATTATTATACACAGAAGAAGAAAGCAATTTGATTGATTCACCGGTACAGCTT 305  
2309 tactagactcttccagactagtgaaagagaatcagaatcttcttggtttacaaagctaa 2368  
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2369 cagaagaagaatggaatataatgaatgatttgaatcttgggaagaggaataatcagaataac 2428  
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2429 tcttaacagcgtttcttcgtcccttcaactctacattctactgctttagaatttaa 2488  
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QY 2489 aaatgacatcttccagaaatgataagatcatatcttgaaattttatataatgta 2548  
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QY 2549 tggaaattcttaggattttttaccagcttggtttaagacccaatgtaaatataaa 2608  
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DB 4 ATA 2

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VERSION BG720949.1 GI:14000136  
KEYWORDS EST.  
SOURCE human.  
ORGANISM Homo sapiens  
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
Mammalia; Eutheria; Primates; Catarrhini; Homnidae; Homo.  
1 (bases 1 to 1019)  
NIH-MGC http://mgi.mc.nih.gov/  
National Institutes of Health, Mammalian Gene Collection (MGC)  
Unpublished (1999)  
Contact: Robert Strausberg, Ph.D.  
Email: egabs-remail.nih.gov  
Tissue Procurement: MIKLOS Palakovits, M.D., Ph.D.  
CDNA Library Preparation: Michael J. Brownstein (NHGRI), Shitaki  
Toshiyuki and Piero Carninci (RIKEN)  
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (ILNL)  
DNA sequencing by: Incyte Genomics, Inc.  
Clone distribution: MGC clone distribution information can be  
found through the I.M.A.G.E. Consortium/ILNL at:  
http://image.llnl.gov  
Plate: LLM10737 row: 1 column: 03  
High quality sequence stop: 785.  
Location/Qualifiers

REFERENCE 1 (bases 1 to 1019)  
AUTHORS NIH-MGC http://mgi.mc.nih.gov/  
TITLES National Institutes of Health, Mammalian Gene Collection (MGC)  
JOURNAL Unpublished (1999)  
COMMENT Contact: Robert Strausberg, Ph.D.  
Email: egabs-remail.nih.gov  
Tissue Procurement: MIKLOS Palakovits, M.D., Ph.D.  
CDNA Library Preparation: Michael J. Brownstein (NHGRI), Shitaki  
Toshiyuki and Piero Carninci (RIKEN)  
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (ILNL)  
DNA sequencing by: Incyte Genomics, Inc.  
Clone distribution: MGC clone distribution information can be  
found through the I.M.A.G.E. Consortium/ILNL at:  
http://image.llnl.gov  
Plate: LLM10737 row: 1 column: 03  
High quality sequence stop: 785.  
Location/Qualifiers

FEATURES  
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BASE COUNT 320 a 221 c 248 g 230 t  
ORIGIN

Query Match 27.7%; Score 753.4; DB 11; Length 1019;  
Best Local Similarity 96.3%; Pred. No. 2.2e-112;  
Matches 814; Conservative 0; Mismatches 26; Indels 5; Gaps 4;

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QY 397 ccgtatgctgaggaagagagcgcgctctgtaacagagcgagggaagcgctgct 456  
DB 65 CCGTAGCTGAGGGAAGAGAGAGCGCGCTCTCTGTCAACAGACCGGGGAAGCGGTGCT 124

QY 457 ttccgagctgcccggtgacacattctccgagaccacatgtagtgcggggagctgc 516  
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 Db 125 TTCCGGGCTGCCCGGTGGACACTTTCCTCGGACCCAGATGATGCGGGCACTGC 184  
 QY 517 catgaactccgagcagatgagatccacagtaaaagacattccagggtgagatccaagt 576  
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 Db 185 CATGAACCTCCGAGCATGAGGATTCACAGTAAGACATTTCGA -GGTGAATCCAACT 243  
 QY 577 caaaaatgaaaaaagacagacatctctgaaatctctgaaactgtaaacaggccagaaaa 636  
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 Db 244 CAAAATGAAAAAACAAGACCATCTGAAATCTGAAAACTGATTAACAGGCCAGAAAA 303  
 QY 637 attcaaatgtaagcactctgggaaagatttctactctgactctctgtaacct 696  
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 Db 304 ATCCAAATGTAAGCCACCTTGGGGAAAAAGTATTTTACCTTGACTTACCTTCGTCAACCAT 363  
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 Db 364 ATCTCAAAAACCTTCAAAAAGACATTAAAGATCTGGGAGGCGCAAGTTGAAGATTCTCAG 423  
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 Db 424 CAAGATATTCAGTATCTTATTTCAAATAGACGGAAGCTAAATTTGCCAACAACCTTG 483  
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 Db 544 CATCCACACCATGATGATGAGATTTCATTTAAGTCACACAGACAGCTGTGTTTAAAGCAGAGA 603  
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 Db 724 TGACAAAAAGAAAAAGAGTGTATTTACTCAAGAAATCAAGTACTTTTCAGTAGAGCAATG 783  
 QY 1113 ggggcaaaagagttgtagtggtagtgcacaaaaaacaagaacaggaagactcaaaaagcctt 1172  
 |||||||  
 Db 784 GGGGCAAAAAGAGTGTGTGTGTGGCAAAAACACAGCACCGGAGAACTCAAGAGAGCCTT 843  
 QY 1173 ttgta 1177  
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 Db 844 GGGTA 848

RESULT 6  
 LOCUS AL559244 725 bp mRNA EST 16-FEB-2001  
 DEFINITION AL559244 LTI\_NFL008\_TC2 Homo sapiens cDNA clone CS0D012YL06 5  
 prime, mRNA sequence.  
 ACCESSION AL559244  
 VERSION AL559244.1 GI:12904555  
 KEYWORDS EST.  
 SOURCE human.  
 ORGANISM Homo sapiens  
 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.  
 REFERENCE 1 (bases 1 to 725)  
 AUTHORS Li, W. B., Gruber, C., Jesse, J. and Polayes, D.  
 TITLE Full-length cDNA libraries and normalization  
 JOURNAL Unpublished (2001)  
 COMMENT Contact: Genoscope  
 Genoscope - Centre National de Sequencage  
 BP 191 91006 Evry cedex - France

FEATURES  
 Source  
 1..725  
 /organism="Homo sapiens"  
 /db\_xref="taxon:9606"  
 /clone="CS0D012YL06"  
 /clone\_1lb="LTI\_NFL008\_TC2"  
 /sex="male"  
 /ruse\_type="T" cells from T cell leukemia"  
 /note="Vector: pCMVSPORT 6; Site 1: NotI; 1st strand cDNA was primed with a NotI-Oligo(dT) primer. Five prime end enriched, double-stranded cDNA was digested with Not I and cloned into the Not I and Eco RV sites of the pCMVSPORT 6 vector. Library was normalized. Library was constructed by Life Technologies. Contact: Feng Liang Life Technologies, Rockville, Maryland 20850, USA Fax: (1) 301 610 8371  
 Email: fliang@lifetech.com URL: http://fulllength.invitrogen.com"

BASE COUNT 224 a 166 c 164 g 171 t  
 ORIGIN

Query Match 26.6%; Score 723.4; DB 10; Length 725;  
 Best Local Similarity 99.9%; Pred. No. 1.6e-107;  
 Matches 724; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 329 cgcgccttgagccgagatccgcgcccgcgaacccgcagctcgcagagcggttaacctactg 388  
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 Db 1 CGGCGCTTGAGGCGGATGTCGGGCGCCGGAACCCGACCTGACAGCGGACTTACTAGC 60  
 QY 389 cgtagaagccgtagctgtgcggaagaagaagcgcgcgtctgtaacagggccggggaa 448  
 |||||||  
 Db 61 CGTAGAGGCGGTAGCTGCGGGAAGAGAGAGGCGGCGTCTGTCAACAGCGCGGGGAA 120  
 QY 449 gcgcgtcttcgcggtgtgcgcggtgagacattctccgagaccagcatgtagtgcgg 508  
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 Db 121 GCGGTGCTTTCGGGCGTCCCGGCGGACACTTCTCCGACCCACATCTAGTGTCCGG 180  
 QY 509 gcgactgcatagaactccgagccatgagatccacagtaaaagacattccaggggtgaa 568  
 |||||||  
 Db 181 GGCAGTGCATGAATCCGAGCCATGAGGATCCACAGTAAGACATTTCCAGGGGTGA 240  
 QY 569 atccaaagtcaaaaaaagaaacagacacatctctgaatctctgaaactgatacagg 628  
 |||||||  
 Db 241 ATCCAAAGTCAAAAATAAAAAACAACACATCTCTGAAACTGTATTAACAGG 300  
 QY 629 ccagaaaatccaaatgtaagccacttgggaaagatttctacttgacttaacctct 688  
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 Db 301 CCAGAAAAATCCAAATGTAAAGCCACTTTGGGCAAAAAGTATTTACTTGAATTAACCTTCT 360  
 QY 689 gtacacatctgaaaaaacttcaaaagacatlaagatctggagggcgagttgaaga 748  
 |||||||  
 Db 361 GTCACCATATTCGAAAAAAGCTTCAAAAGACATTAAGATCTGGAGGGGAGTTGAAGAA 420  
 QY 749 ttctcagaagaagatcagttatcttatttcaataaagaagaactaaattgcaaa 808  
 |||||||  
 Db 421 TTTCTGAGCAAAATATGATGATTTATTTTCAAAAGAGAAAGAACTTAATTTGCACAA 480  
 QY 809 acctgtgcgaattctctctgacccaagctccgaatctgcacatactgaagaacct 868  
 |||||||  
 Db 481 ACCTTGCGTCGAATTTCTCTGTACCAAGTCCGAATTAATTAATTAATTAATTAATTAAT 540  
 QY 869 taacctatccagcagcagatggaagttcaatlaagtcacagacagatglttgaagc 928  
 |||||||  
 Db 541 TCACCTCATCCCGACCATATGATGAATTTAATTAATTAATTAATTAATTAATTAATTAAT 600  
 QY 929 agaggaataatttagttgaagaagatcaagagacatgatttcttcttaaatagc 988  
 |||||||  
 Db 601 AGAGGAAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAAT 660  
 QY 989 atattcaaatgctgtcagtgaggaagaaatcttcatattgatacatgataac 1048  
 |||||||

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Db      661 AATATATCAATGCTTGTGTCAGGGAGTAAATCTTCATATGATGACATTAGATAC 720
OY      1049 tacat 1053
        |||||
Db      721 TACAT 725

RESULT 7
AUI28881 903 bp mRNA EST 24-OCT-2000
LOCUS AUI28881 NT2RP2 Homo sapiens cDNA clone NT2RP2004396 5', mRNA
DEFINITION
ACCESSION AUI28881.1 GI:10989235
VERSION AUI28881.1
KEYWORDS EST.
SOURCE human.
ORGANISM Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Homnidae; Homo.
1 (bases 1 to 903)
Ota,T., Nishikawa,Y., Suzuki,Y., Ishii,S., Saito,K., Kawai,Y.,
Yamamoto,J., Nakamatsu,A., Nakamura,Y., Nagai,T., Sugano,S. and
Ishigai,T.
HRI human cDNA project
Unpublished (2000)
JOURNAL
COMMENT
CONTACT: Takao Isegai
Genomics Laboratory
153-3 Yata, Katsushika-ku, Chiba 292-0812, Japan
Tel.: 81-438-543951
Fax: 81-438-52-3952
Email: genomics@hri.co.jp
HRI human cDNA project, 5' - 3'-end one pass sequencing; Helix
Research Institute, cDNA library construction; Department of
Virology, Institute of Medical Science, University of Tokyo, and
Helix Research Institute
FEATURES
Source
1..903 /Qualifiers
/organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="NT2RP2004396"
/clone_1lb="NT2RP2"
/clone_1lb="hepatocarcinoma"
/cell_type="NT2"
/vector="PMI18FRL3; mRNA from NT2 neuronal precursor
cells after 2-weeks retinoic acid (RA) induction;"
BASE COUNT 326 a 165 c 157 g 242 t 13 others
ORIGIN
Query Match 26.5% Score 721.6; DB 10; Length 903;
Best Local Similarity 92.3%; Pred. No. 3e-107;
Matches 820; Conservative 0; Mismatches 56; Indels 12; Gaps 6;
OY 1593 taatttcgaagaagtcggaagaatgatacaagc-agaagaagaaattcctgt 1651
Db 1 TATTTCTCAAAAAGTTTCAGGAAGATGATACACAGCGAAGACAGAAATTCTCTG 60
OY 1652 aataaagacccaggaagaactgaataaaagctctgttttttcagaagccatcccccacc 1711
Db 61 ATTAAGAGCCACGAGAACTGAAAAAGCTCTGTTTATTTTCACAGCCCATCCCCACC 120
OY 1712 ctccaataaattgagaaggttaataagaagaattgttaataatgttcacgttaagta 1771
Db 121 CTCCTAATGAATTGAGCGCTTAAATGAGAAATGATTAATGTTCCATGTTAAGTA 180
OY 1772 cagctgaagaatacagaagaatttcacagctacactacataaataaacaagaag 1831
Db 181 CACCTGAAGTGCATTAAGACGAATTTTACACAGCTACCTACATFAAANAACAGG 240
OY 1832 aatgcatcttcaacatttcgaacacacatttaagtaagaactagaagaagaag 1891
Db 241 AATGCATCTCTGACATTTCGAGACACACATTAAATGTAATGACTTACAGACTANGG 300

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OY 1892 taagtaactaataatgatacatagagcatctgtacatgtttctgatttcagtaagata 1951
Db 301 TAGATCACTATTAATGTAAATGTAACATACAGGCATCTGTACATGTGTTCTGATTCAGTACAGATA 360
OY 1952 atagtgatctcgaacacaaagaagtcagatactgtgtttttccacaagaagatctca 2011
Db 361 ATAGTGATCTCACCAAAACAGAAAGTCAGATCTGTGCTTTTCCACAAGAGATCTCA 420
OY 2012 aggaagaagacccatctcaatatttactcatgattctgtctgtgataaacaataggtc 2071
Db 421 AGGAAAAGACCTCTTCAATATTTCATCATGATTCGTGCTGTGATTAACAATACAGTT 480
OY 2072 cacaagaagacccatctgtctcagaagaagccatctcactcctcgtgaagagacc 2131
Db 481 CACAAGACACCTTAACCTGTCAGCAGCAGCCATTCATCTCCTCTGAGCAGACCCA 540
OY 2132 atgaatgtactcaagaataatgatagttactcctcgtgtaaatatatacagaagaatga 2191
Db 541 ATGAATGTGACTTCAAGAAATATGATATGATTTACCTCTGTGTAATAATACATCGAANAAGTGA 600
OY 2192 aataaatatta-aggagaataagaagaataatctggaacacaaatgctgaattgataaa 2250
Db 601 AATTAATATTAGGACGAATATGAAAAGAAAATCTGGAAACCAAAATCTGAATTTGATPAA 660
OY 2251 agaatgaatttattacacaagaagaagaagaagaattgtgtagttcacacgggtacagttcta 2310
Db 661 AGAAGCTGAATTTATTTNCCCAAGACAAACNGAATTTGT-TTTCACCGGTACAGCTTTTA 719
OY 2311 ctgaactgttccaagctagtgaaagaagaacagaatttttggttttcacaagctacaca 2370
Db 720 CTANAAGCTGTTACAGCTAGTACGAAAAAATCNAATTTTGGCTTTCACAAAGCTACCCA 779
OY 2371 gaaa--gagtgatcatgcaatgttttagatattgg----aagaggaataatcaga 2423
Db 780 GAAAAAATAATTGGTATATGCCATGTGTTTANATATATGGGGGAAAAAATTTCCNAAT 839
OY 2424 taactgttaacagcgtttt--ctcgccccctcactacatc 2469
Db 840 AATCGTGTAAACAGCGTTTTCGCCNGTCCTTCAACTTCCTCCWTT 887

RESULT 8
AL580899/c 742 bp mRNA EST 16-FEB-2001
LOCUS AL580899 LRI_NFL008_Tc2 Homo sapiens cDNA clone CS0DJ012YL06 3
DEFINITION
ACCESSION AL580899
VERSION AL580899.1 GI:12947367
KEYWORDS EST.
SOURCE human.
ORGANISM Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Homnidae; Homo.
REFERENCE 1 (bases 1 to 742)
AUTHORS Li,W.B., Gruber,C., Jessee,J. and Polayes,D.
TITLE Full-length cDNA libraries and normalization
JOURNAL Unpublished (2001)
COMMENT Contact: Genoscope
Genoscope - Centre National de Sequencage
BP 191 91006 EVRY cedex - France
Email: seqref@genoscope.cns.fr, web : www.genoscope.cns.fr.
FEATURES
Source
1..742
/organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="CS0DJ012YL06"
/clone_1lb="LRI_NFL008_Tc2"
/sex="male"
/tissue_type="T cells from T cell leukemia"
/note="Vector: PCWVSPORT 6; Site:1; NotI; 1st strand cDNA
was primed with a NotI-oligo(dT) primer. Five prime
end enriched, double-stranded cDNA was digested with Not I and

```

cloned into the Not I and Eco RV sites of the pCMVSPORT 6 vector. Library was normalized. Library was constructed by Life Technologies. Contact : Feng Liang Life Technologies, a division of Invitrogen 9600 Medical Center Drive Rockville, Maryland 20850, USA Fax : (1) 301 610 8371 Email : fliang@lifetech.com URL : <http://fulllength.invitrogen.com>

BASE COUNT 228 a 124 c 120 g 269 t 1 others

Query Match 26.4%; Score 719; DB 10; Length 742;

Best Local Similarity 99.1%; Pred. No. 8.3e-107;

Matches 734; Conservative 0; Mismatches 5; Indels 2; Gaps 1;

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QY 1912 atcaagcatctgtacatgttctgtattcagtlacagataatagtgtatctcaacaa 1971
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DB 742 ATACAGCATCTGTACATGTTCTGATTTTCAGTACAGATTAATAGTGTCAACCAAA 683
    |||||||
QY 1972 cagaagtcagatctgtgtctttccagcaagatctcaaggaaggaagacctattca 2031
    |||||||
DB 682 CAGAAGTCAGATCTGTGCTTTTTCACCAAGATCTCAAGGAAGGACCTTCAATCA 623
    |||||||
QY 2032 atatttactatgtatctgtctgtatcaaatcaacagttcaacagacactaactgt 2091
    |||||||
DB 622 ATATTACTCATGATCTGTGCTGTATTAACATTAACATTCACAGACACCTAACTGT 563
    |||||||
QY 2092 caggcaagctccatctcactactcctctcgaggaaacccaatgaatgttacttcaaga 2151
    |||||||
DB 562 CAGGCAAGGCTCCATCTCCATCTCTCTGAGAACCAATGATGTACTTCAAGAAAT 503
    |||||||
QY 2152 atgataatcttaccctctgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgt 2211
    |||||||
DB 502 ATGGTATCTTACCTCTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGT 443
    |||||||
QY 2212 agaaagaagaatctcgaaacccaatctgtcaattgtataaagacgtcaatttatacaca 2271
    |||||||
DB 442 AGAAAGAAGAATCTGGAACCAATCTGATTTGTATAAGACGAAATTTTATACACA 383
    |||||||
QY 2272 gaagaaacagaattgttagttacacgggtacagttcttactagactgtttcagaactgt 2331
    |||||||
DB 382 GAAGAAACAGAAATTTGTAGTTCACCGGTACAGTCTTACTGACTTGTTCAGACTAGT 323
    |||||||
QY 2332 gaagagaagaatcaatttgggtttcacaagctcacagaaagaagtgatgtacat 2391
    |||||||
DB 322 GAAGAGAATGAGAAATTTTGGTTTCAACAGCTACAGAAAGAGTGATTAATGCAAT 263
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QY 2392 gttttagatatttgggaaggaagaatcag--ataatcgttcaacagcgttttctcgt 2449
    |||||||
DB 262 GTTTAGATATTTGGGAAGAGGAATAATTCAGCATACATCTGTTAACAGGTTTCTCGT 203
    |||||||
QY 2450 ccccttcaactctacattactgtgcttttagaatttaaaaaatgacatttccagaag 2509
    |||||||
DB 202 CCCCTTCAACTTCTCATATGATGACTGGCTTTTAGAAATCTAAAAATCATCTTTTGAAG 143
    |||||||
QY 2510 tgataagaataatcttctgtgaattttataaatagtatgtgaatctttagattttt 2569
    |||||||
DB 142 TGATTAAGATCATATTTGTAATTTTATTAATATGTATGAAATTCCTAGATTTT 83
    |||||||
QY 2570 taccagcttctgttacaacccaatgtlaatatataaataaataatgtcaatttcta 2629
    |||||||
DB 82 TACACGCTTTGTTTACACACCAATGTAAATATTAATAATTTTGCATTTTCTA 23
    |||||||
QY 2630 cagaatgtaatcctgttaaa 2650
    |||||||
DB 22 CAGAAATGAAATACCTGTAA 2
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RESULT 9  
 A1948485/c A1948485 744 bp mRNA EST 08-MAR-2000  
 LOCUS wq6d08.x1 NCI-CGAP\_K1d12 Homo sapiens cDNA clone IMAGE:2470479 3'  
 DEFINITION similar to TR:075226 075226 WUGSC:H\_RGI135C18.1 PROTEIN ;, mRNA

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sequence.
ACCESSION A1948485
VERSION A1948485.1 GI:5740795
KEYWORDS EST.
SOURCE human.
ORGANISM Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Homnidae; Homo.
REFERENCE 1 (bases 1 to 744)
AUTHORS NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
TITLE National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
JOURNAL Tumor Gene Index
COMMENT Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
Email: cgap@remail.nih.gov
Tissue Procurement: Christopher Moskaluk, M.D., Ph.D., Michael R.
Emmert-Buck, M.D., Ph.D.
CDNA Library Preparation: M. Bento Soares, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
www.bio.lnl.gov/db/brp/image/image.html
Insert length: 1484 Std Error: 0.00
Seq primer: -40up from gibco
High quality sequence stop: 432.
FEATURES
Location/Qualifiers
source 1..744
/organism="Homo sapiens"
/db_xref="taxon:9606"
/clone_image="IMAGE:2470479"
/clone_id="NCI-CGAP_K1d12"
/tissue_type="2 pooled tumors (clear cell type)"
/lab_host="DH10B"
/note="Organ: Kidney; Vector: pT73D-Pac (pharmacia) with
a modified polylinker; Site: 1: Not I; Site 2: Eco RI;
Plasmid DNA from the normalized library NCI-CGAP_K1d5 was
prepared, and as circles were made in vitro. Following HAP
purification, this DNA was used as tracer in a subtractive
hybridization reaction. The driver was PCR-amplified cDNAs
from a pool of 5,000 clones made from the same library
(cloneids 1323912-1325831, 1471368-1472903 and
1492104-1493255). Subtraction by Bento Soares and M.
Fatima Bonaldo."
BASE COUNT 223 a 127 c 117 g 277 t
ORIGIN
Query Match 26.1%; Score 710; DB 10; Length 744;
Best Local Similarity 97.3%; Pred. No. 2.4e-105;
Matches 722; Conservative 0; Mismatches 20; Indels 0; Gaps 0;
QY 1900 tataatgttaacatacagcactgtacatgttctgtattcagtlacagataatagtga 1959
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DB 744 TATCATATGTAAACACAGCGTCGTGATCATGTTTGTGATCTCAGTACATCATAGTGA 685
    |||||||
QY 1960 tctcaaccaaacagaagtcagatctgtcttttccagcaagagatctcaaggaag 2019
    |||||||
DB 684 TCTCAACCGAGACAGAGTCAGATATCTGTGCTTTTCCAGCAAGGATCTCAAGGAACG 625
    |||||||
QY 2020 gaccttcatcaatatttctcattgttctgtctgtataaataaaggttcaacaaag 2079
    |||||||
DB 624 GACCTTCATTCGATTTTCTCATGTATCTGGTCTGATTAACATTAACAGTTTCAACAGAG 565
    |||||||
QY 2080 caactaacgttcaaggcaaggctccatctcactcctctgaggaacccaatgaatgt 2139
    |||||||
DB 564 CACTTAACGTGTTCAAGCAAGGCTCATTCATTCCTCTGAGAGAACCATGAATGT 505
    |||||||
QY 2140 gacttcaagaataatgatagttaccttctgtgtataaataacatcgaagaagtgaataata 2199
    |||||||
DB 504 GACTTCMAAGAAATATGATAGTTTACCTCTGTGTAATAATCATTCGAAAGTGAATAATAA 445
    |||||||
QY 2200 ttaggacgaataatagaataaataatctcgaaacaaatgctggaatttgataaagaactgaa 2259
    |||||||

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Db 444 TTAGAGAAATGAAAGAAATCTGAGAACCAATCTGAATTTGATAAAGAACGTAA 385
    |||
Oy 2260 ttatacacaagaagaagaacgaattcttagtcaccggtagacgtcttactagactg 2319
    |||
Db 384 TTATATACACAGAGAGAAACGAATTTGATGATTCACCGGACAGCTTACTAGACTTG 325
    |||
Oy 2320 tttagcagctagtagaagaagaacagaattttgggtttcacaagctcacaagaagaagat 2379
    |||
Db 324 TTTCAGACTGCTGACAGAGAAATCAGAAATTTTGGCTTTCACAGCTACAGAGAAAGAGT 265
    |||
Oy 2380 ggtatgatgaatgtttatagatattgggaagaagaagaatcagaatactgtaaacag 2439
    |||
Db 264 GGTATATGCAATGTTTATGATTTGGGAGAGAGAAATTCAGTATATCTGTTACAGCG 205
    |||
Oy 2440 tttcttcgtccctcacaactcacaatttactctggtctttagaatttaaaatgcatc 2439
    |||
Db 204 TTTTCTGTCGTCCTTCACTCTGACATTTACTGCTGCTTTAGAAATTAATAAATGCTATC 145
    |||
Oy 2500 ttctcagaagtgtatgaagatcatctctgaaattcttataatagatgaaattctt 2559
    |||
Db 144 TTTTCAGAGAGTGAATGAAGATCATATTTCTTGAAATTTTATATGATANGAAATCTT 85
    |||
Oy 2560 aggatcttttaccagctgtgttaccagaccacaatgtaaatltaaaataatattg 2619
    |||
Db 84 AGATTTTATTTACAGCTTGTGTTACAGCCAAATGTAATTTAAATTAATATTTTG 25
    |||
Oy 2620 caattctcagaatgata 2641
    |||
Db 24 AAAAAATCTTAAAAA 3
    |||
RESULT 10
Bg496289 821 bp mRNA EST 27-MAR-2001
LOCUS 602538259F1 NIH_MGC_59 Homo sapiens cDNA IMAGE:465946 5'
DEFINITION mRNA sequence.
ACCESSION Bg496289
VERSION Bg496289.1 GI:13457805
KEYWORDS EST.
SOURCE human.
ORGANISM Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE 1 (bases 1 to 821)
AUTHORS NIH-MGC http://imgc.ncl.nih.gov/
TITLE National Institutes of Health, Mammalian Gene Collection (MGC)
JOURNAL Unpublished (1999)
COMMENT Contact: Robert Strausberg, Ph.D.
Email: rgs@bbs-femail.nih.gov
Tissue Procurement: AMCC
CDNA Library Preparation: CLONTECH Laboratories, Inc.
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Invitrogen Genomics, Inc.
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: L10M435 row: e column: 15
High quality sequence stop: 742.
Location/Qualifiers
1. 821
/organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:465946"
/clone_lib="NIH_MGC_59"
/tissue_type="muscleepidermoid carcinoma"
/lab_host="DH10B (TI phase-resistant)"
/notes="Organ: Lung; Vector: pDR-LIB (Clontech); Site: 1:
SfiI (ggcgctcgagc); Site: 2: SfiI (ggcattatggc);
Double-stranded cDNA was prepared from cell line RNA. 5'
and 3' adaptors were used in cloning as follows: 5'
adaptor sequence: 5'-CACGGCCATTTAGGCC-3' and 3' adaptor
sequence: 5'-ATTCTAGAGCCAGCGGCCGACATG-dT(30)BN-3'

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BASE COUNT 273 a 166 c 176 g 206 t
ORIGIN
Query Match 25.6%; Score 696.4; DB 11; Length 821;
Best Local Similarity 97.2%; Pred. No. 3,7e-103;
Matches 762; Conservative 0; Mismatches 16; Indels 6; Gaps 5;
    (where B = A, C, or G and N = A, C, G, or T). Average
    insert size 1.65 kb (range 0.9-4.0 kb). 15/15 clones
    contained inserts by PCR. This library was enriched for
    full-length clones and was constructed by Clontech
    Laboratories (Palo Alto, CA). Note: this is a NIH-MGC
    library."
Oy 417 gaagggcgccgctctgtcacaagcgagggaagccgtgtcttgaggtgctgcccgttgga 476
    |||
Db 1 GAGCGGCGCTCTGTCACACAGCGCGGAGAGCCGTGCTTTCGCGCTGCCCGGTGCGA 60
    |||
Oy 477 caatttcgccgagccagcagatgtaggtgcggcgactgcattgaactccggagccatga 536
    |||
Db 61 CACTTTCGCGACCCAGCATGTAGTGTCCGGGCGACTGCCATGAATCTCGGAGCATGA 120
    |||
Oy 537 ggalccacagtaagaagacatctccagggtggaatccaaagtcgaaatgaaatgaaacagac 596
    |||
Db 121 GGAATCCAGTAAGGACATTTCCAGGGTGGAATCCAAAGTCAAAAATGAAAAACAGAC 180
    |||
Oy 597 catctcgaatctctgaaactga taacagggccagaaaaatccaaatgtgaagccactt 656
    |||
Db 181 CATCTGGAATCTGTGAAACTGATAACAGGCCAGAAAAATCAATGTAGACCATTT 240
    |||
Oy 657 ggggaagaatgtttacacttgaacttaacttctgacacatatctgaaatcacaag 716
    |||
Db 241 GGGGAAAAGATTTTACCTTGACTTACTTCTGTGACATATGAAAAATCTTCAAAAG 300
    |||
Oy 717 acattaagatctgggaaggcgagttgaagaattctcacaagaatcagatcacta 776
    |||
Db 301 ACATTAAAGATCTGGGAGGGGCGAGTGAAGATTTCTCACAAAGATACATTATCTTA 360
    |||
Oy 777 ttcaaatgaagaagaagactaaatttcacaacaccttggtgcgaattctctctgaacaa 836
    |||
Db 361 TTTCAAAATGAAGAGAGCTAAATTTGCAAAACCTTGAGTGAATTTCTCTGTACCA 420
    |||
Oy 837 gtccagaatctgata tactgcagaacacctcacctcattccagcagatgtagaagt 896
    |||
Db 421 GTCCAGATCTGCATATCTGACAGAAACCTTCACTCATTCACGCAATGAGAGATT 480
    |||
Oy 897 catlaagtcacagacacagtggttttaagcagaaggaataatcagtagtgaagaagcta 956
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Db 481 CATTTAAGTCACACAGACAGTGTGTTAAAGCAGAGAAATTAATTAAGTTGAAAAAGCTA 540
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[illegible]

JOURNAL COMMENT  
 Unpublished (1999)  
 Contact: Robert Strausberg, Ph.D.  
 Email: cghps-femail.nih.gov  
 Tissue Procurement: ATCC  
 cDNA Library Preparation: Life Technologies, Inc.  
 DNA Sequencing by: Incyte Genomics, Inc.  
 Clone distribution: MGC clone distribution information can be  
 found through the I.M.A.G.E. Consortium/LLNL at:  
 http://image.llnl.gov  
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 Note: this is a NH-MGC Library."  
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AUTHORS	Hegde,P., Qi,R., Abernathy,K., Dharrap,S., Gasparé,R., Gay,C., Holtz,I.E., Saeed,A.I., Sharov,V., Lee,N.H., Yeatman,T.J. and Quackenbush,J.		
TITLE	Assessment of gene expression patterns in a model of colon tumor metastasis using a 19,200 element cDNA microarray		
JOURNAL	Unpublished (2000)		
COMMENT	Contact: John Quackenbush The Institute for Genomic Research 9712 Medical Center Dr., Rockville, MD 20850, USA Tel: 301 838 3528 Fax: 301 838 0208 Email: john@etigr.org Plate: 150		
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VERSION	A1761101.1		
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TITLE	Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.		
JOURNAL	1 (bases 1 to 743)		
COMMENT	NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.		
	National Cancer Institute, Cancer Genome Anatomy Project (CGAP),		
	Tumor Gene Index		
	Unpublished (1997)		
	Contact: Robert Strausberg, Ph.D.		
	Email: cgaabs@femail.nih.gov		
	Tissue Procurement: Christopher Moskaluk, M.D., Ph.D., Michael R.		
	Emmert-Buck, M.D., Ph.D.		
	CDNA Library Preparation: M. Bento Soares, Ph.D.		
	CDNA Library Arrayed by: Greg Lennon, Ph.D.		
	DNA Sequencing by: Washington University Genome Sequencing Center		
	Clone distribution: NCI-CGAP clone distribution information can be		
	found through the I.M.A.G.E. Consortium/ILMUT at:		
	www-bio.illn.gov/bbrp/image/image.html		
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	hybridization reaction. The driver was PCR-amplified cDNAs		
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	(cloneids 1323912-1325831, 1471366-1472903 and		
	1492104-1493255). Subtraction by Bento Soares and M.		

Fri Dec 28 08:22:48 2001

us-09-830-647-4.rst

Page 12

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ORIGIN Fatima Bonaldo, \*

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